













qb/Aa548636/AA548636 nj 38d02.si NCI CGAP AA1 Homo Sapiens cDNA clone lwaGE 994755 similar to gb:246633 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-2 a*0201 ALPHA (HUMAN); qb/R12066/R12066 yf54d10.r1 Homo sapiens cDNA clone 26062 5'similar to qb:X00492_cds 1 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-3A*0301

- +2 -05, p = 8.1e-05 = 18/18 (100%), Frame Score= 30 (41.9 bits), Expect = 8.1e-05,

18 VDDTQEVRFDSDAASQRM -

152

Sbjct:

Length = 581

gb/AA132653/AA132653 zo2la05.rl Stratagene colon (#937204) Homo sapiens cDNA clone 587504 5' similar to gb:z46633 HLA CLASS I HISTOCCMPATIBILITY ANTIGEN, A-2 A* 0201 ALPHA (HUMAN);

211

158 VDDTQFVRFDSDAASQRM

1 VDDTQFVRFDSDAASQRM 18

Query: Sbjct:

Plus Strand HSPs:

alpha (human);

Length = 405

VDDTQFVRFDSDAASQRM

13 VDDTQFVRFDSDAASQRM 66

qb/R59764/R59764 yh07c05.r1 Homo sapiens cDNA clone 42563 5'similar to qb:X00492 cds1 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-3 A* 0301 ALPHA (HUMAN);.

VDDTQFVRFDSDAASQRM 203

Sbjct: 158

VDDTQEVREDSDAASQRM

Query: 1 VDDTQFVRFDSDAASQRM 18

Score = 90 (41.9 bits), Expect = 6.1e-05, P = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame =+1

Plus Strands HSPs:

Length = 461

1 VDDTQFVRFDSDAASQRM 18 Query:

Sbjct: 154 VDDTQFVRFDSDAASQRM 207

19<u>04/R13904</u> yf62c03.rl Homo sapiens cDNA clone 26801 5'similar t gb:m64742_cds1 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-23(A-9) qb/R13904/R13904 yf62c03.r1 Homo A*2301 (HUMAN);

qb/AA488534/AA488534 ab37f08.rl Stratagene Hela cell s3 937216 Homo sapiens cDNA clone 843015 5' similar to gb:L06425 HLA CLASS I HISTOCCMPATIBILITY ANTIGEN, AW-34(A-10)) A*3402 ALPHA (HUMAN):

138 VDDTQFVRFDSDAASQRM 210

VDDTQFVRFDSDAASQRM

Query: 1 VDDTQFVRFDSDAASQRM 18

ć

ເນ ::

ಭ

Score 90 (41.9 bits), Expect = 8.1e-05 P=8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame = +2

£

Frame

(41.9 bits), Expect = 8.1e-05, P = 8.1e-05 = 18/18 (100%), Positives = 18/18 (100%),

Score = 90 Identities

Strand HSPs:

Plus

Length = 478

VDDTQFVRFDSDAASQRM

Query: Sbjct:

...

VDDTQEVRFDSDAASQRM Sbjct: 146 VDDTQEVRFDSDAASQRM 199

Identities = 18/18 (100%), Positives Plus Strand HSPs: Length -Score = 90 (41.9 bits), Expect = 8.1e-05, P=9.1e-05 Identities = 18/18 (100%), Positives=18/18(100%), Frame = +2

VDDTQFVRFDSDAASQRM VDDTQFVRFDSDAASQRM 205 Query:

qb/Aa147151/Aa147151 zo32d06.rl Stratagene colon (#937204) Homo sapiens cDNA clone 5885875 similar to gb:M64740 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-24 (A-9) A*2402 ALPHA (HUMAN);

Plus Strands HSPs:

Score = 90 (41.9) bits), Expect = 8.1e-05, P = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame

£ =

Score = 90 (41.9) bits), Expect = 8.1e-05,P=8.1e-05 Identities = 18/18 (100%) Positives=18/18(100%) Frame

Plus Strand HSPs:

Length - 428

+

1 VDDTQFVRFDSDAASQRM 18 VDDTQFVRFDSDAASQRM Query:

gb/ H23377/H23377 ym57e02.rl Homo sapiens cDNA clone 52227 5'similar to gb:x00492 cds1 HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, A-3A A*0301 ALPHA (HUMAN);

Length = 459

Strand HSPs: Plus Score = 90 (41.9 bits), Expect = 8.1e-05, P = 8.1e-05 Identities = 18/18 (100%), Positives = 18/18 (100%), Frame

7

VDDTQEVREDSDAASQRM

Length = 459 Plus Strand HSPs:

1 VDDTQFVRFDSDAASQRM 18 Query:

